



IFWO

RAW SEQUENCE LISTING

DATE: 07/21/2004

PATENT APPLICATION: US/10/786,774

TIME: 09:27:25

Input Set : A:\08550~1.txt

Output Set: N:\CRF4\07212004\J786774.raw

3 <110> APPLICANT: Holoshitz, Joseph
 4 Ling, Song
 6 <120> TITLE OF INVENTION: Methods and Compositions for the Treatment of Diseases
 Associated
 7 with Signal Transduction Aberrations
 9 <130> FILE REFERENCE: UM-08550
 11 <140> CURRENT APPLICATION NUMBER: 10/786,774
 C--> 12 <141> CURRENT FILING DATE: 2004-02-25
 14 <160> NUMBER OF SEQ ID NOS: 36
 16 <170> SOFTWARE: PatentIn version 3.2
 18 <210> SEQ ID NO: 1
 19 <211> LENGTH: 5
 20 <212> TYPE: PRT
 21 <213> ORGANISM: Artificial Sequence
 23 <220> FEATURE:
 24 <223> OTHER INFORMATION: Synthetic
 26 <400> SEQUENCE: 1
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 29 1 5
 32 <210> SEQ ID NO: 2
 33 <211> LENGTH: 5
 34 <212> TYPE: PRT
 35 <213> ORGANISM: Artificial Sequence
 37 <220> FEATURE:
 38 <223> OTHER INFORMATION: Synthetic
 40 <400> SEQUENCE: 2
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 43 1 5
 46 <210> SEQ ID NO: 3
 47 <211> LENGTH: 5
 48 <212> TYPE: PRT
 49 <213> ORGANISM: Escherichia coli
 52 <220> FEATURE:
 53 <221> NAME/KEY: MISC_FEATURE
 54 <222> LOCATION: (2)..(2)
 55 <223> OTHER INFORMATION: The residue at this position can be lysine or arginine.
 57 <220> FEATURE:
 58 <221> NAME/KEY: MISC_FEATURE
 59 <222> LOCATION: (3)..(4)
 60 <223> OTHER INFORMATION: The residues at these positions can be any amino acid.
 62 <400> SEQUENCE: 3
 W--> 64 Gln Xaa Xaa Xaa Ala
 65 1 5
 68 <210> SEQ ID NO: 4

(pg. 6)

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69 <211> LENGTH: 5
70 <212> TYPE: PRT
71 <213> ORGANISM: Escherichia coli
74 <220> FEATURE:
75 <221> NAME/KEY: MISC_FEATURE
76 <222> LOCATION: (3)..(4)
77 <223> OTHER INFORMATION: The residues at these positions can be any amino acid.
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82 1 5
85 <210> SEQ ID NO: 5
86 <211> LENGTH: 15
87 <212> TYPE: PRT
88 <213> ORGANISM: Artificial Sequence
90 <220> FEATURE:
91 <223> OTHER INFORMATION: Synthetic
93 <400> SEQUENCE: 5
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96 1 5 10 15
99 <210> SEQ ID NO: 6
100 <211> LENGTH: 14
101 <212> TYPE: PRT
102 <213> ORGANISM: Artificial Sequence
104 <220> FEATURE:
105 <223> OTHER INFORMATION: Synthetic
107 <400> SEQUENCE: 6
109 Lys Asp Leu Leu Gln Lys Arg Ala Ala Val Asp Thr Tyr
110 1 5 10
113 <210> SEQ ID NO: 7
114 <211> LENGTH: 15
115 <212> TYPE: PRT
116 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Synthetic
121 <400> SEQUENCE: 7
123 Lys Asp Ile Leu Glu Asp Glu Arg Ala Ala Val Asp Thr Tyr Cys
124 1 5 10 15
127 <210> SEQ ID NO: 8
128 <211> LENGTH: 14
129 <212> TYPE: PRT
130 <213> ORGANISM: Artificial Sequence
132 <220> FEATURE:
133 <223> OTHER INFORMATION: Synthetic
135 <400> SEQUENCE: 8
137 Lys Asp Ile Leu Glu Asp Glu Arg Ala Ala Val Asp Thr Tyr
138 1 5 10
141 <210> SEQ ID NO: 9
142 <211> LENGTH: 15
143 <212> TYPE: PRT

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144 <213> ORGANISM: Artificial Sequence
146 <220> FEATURE:
147 <223> OTHER INFORMATION: Synthetic
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152 1 5 10 15
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156 <211> LENGTH: 15
157 <212> TYPE: PRT
158 <213> ORGANISM: Artificial Sequence
160 <220> FEATURE:
161 <223> OTHER INFORMATION: Synthetic
163 <400> SEQUENCE: 10
165 Lys Asp Leu Leu Glu Gln Arg Arg Ala Ala Val Asp Thr Tyr Cys
166 1 5 10 15
169 <210> SEQ ID NO: 11
170 <211> LENGTH: 5
171 <212> TYPE: PRT
172 <213> ORGANISM: Artificial Sequence
174 <220> FEATURE:
175 <223> OTHER INFORMATION: Synthetic
177 <400> SEQUENCE: 11
179 Gln Lys Arg Leu Ala
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184 <211> LENGTH: 5
185 <212> TYPE: PRT
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189 <223> OTHER INFORMATION: Synthetic
191 <400> SEQUENCE: 12
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197 <210> SEQ ID NO: 13
198 <211> LENGTH: 5
199 <212> TYPE: PRT
200 <213> ORGANISM: Escherichia coli
203 <220> FEATURE:
204 <221> NAME/KEY: MISC_FEATURE
205 <222> LOCATION: (2)..(2)
206 <223> OTHER INFORMATION: The residue at this position is dLys.
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211 1 5
214 <210> SEQ ID NO: 14
215 <211> LENGTH: 5
216 <212> TYPE: PRT
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:

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234 <223> OTHER INFORMATION: Synthetic
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239 1 5
242 <210> SEQ ID NO: 16
243 <211> LENGTH: 5
244 <212> TYPE: PRT
245 <213> ORGANISM: Artificial Sequence
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257 <211> LENGTH: 169
258 <212> TYPE: PRT
259 <213> ORGANISM: Artificial Sequence
261 <220> FEATURE:
262 <223> OTHER INFORMATION: Synthetic
264 <400> SEQUENCE: 17
266 Met Asp Ile Asp Pro Tyr Lys Glu Phe Gly Ala Thr Val Glu Leu Leu
267 1 5 10 15
270 Ser Phe Leu Pro Ser Asp Phe Phe Pro Ser Val Arg Asp Leu Leu Asp
271 20 25 30
274 Asn Ala Ser Ala Leu Tyr Arg Glu Ala Leu Glu Ser Pro Glu His Cys
275 35 40 45
278 Ser Pro His His Thr Ala Leu Arg Gln Ala Ile Leu Cys Trp Gly Glu
279 50 55 60
282 Leu Met Thr Leu Ala Thr Trp Val Gly Gly Asn Leu Glu Asp His Lys
283 65 70 75 80
286 Asp Leu Leu Glu Gln Lys Arg Ala Ala Val Asp Thr Tyr Cys Val Asp
287 85 90 95
290 Pro Ile Ser Arg Asp Leu Val Val Ser Tyr Val Asn Thr Asn Met Gly
291 100 105 110
294 Leu Lys Phe Arg Gln Leu Leu Trp Phe His Ile Ser Cys Leu Thr Phe
295 115 120 125
298 Gly Arg Glu Thr Val Ile Glu Tyr Leu Val Ser Phe Gly Val Trp Ile
299 130 135 140
302 Arg Thr Pro Pro Ala Tyr Arg Pro Pro Asn Ala Pro Ile Leu Ser Thr
303 145 150 155 160

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Input Set : A:\08550~1.txt

Output Set: N:\CRF4\07212004\J786774.raw

306 Leu Pro Ala Trp Ala Arg Val Ile Asn

307 165

310 <210> SEQ ID NO: 18

311 <211> LENGTH: 169

312 <212> TYPE: PRT

313 <213> ORGANISM: Artificial Sequence

315 <220> FEATURE:

316 <223> OTHER INFORMATION: Synthetic

318 <400> SEQUENCE: 18

320 Met Asp Ile Asp Pro Tyr Lys Glu Phe Gly Ala Thr Val Glu Leu Leu

321 1 5 10 15

324 Ser Phe Leu Pro Ser Asp Phe Phe Pro Ser Val Arg Asp Leu Leu Asp

325 20 25 30

328 Asn Ala Ser Ala Leu Tyr Arg Glu Ala Leu Glu Ser Pro Glu His Cys

329 35 40 45

332 Ser Pro His His Thr Ala Leu Arg Gln Ala Ile Leu Cys Trp Gly Glu

333 50 55 60

336 Leu Met Thr Leu Ala Thr Trp Val Gly Gly Asn Leu Glu Asp His Lys

337 65 70 75 80

340 Asp Ile Leu Glu Asp Glu Arg Ala Ala Val Asp Thr Tyr Cys Val Asp

341 85 90 95

344 Pro Ile Ser Arg Asp Leu Val Val Ser Tyr Val Asn Thr Asn Met Gly

345 100 105 110

348 Leu Lys Phe Arg Gln Leu Leu Trp Phe His Ile Ser Cys Leu Thr Phe

349 115 120 125

352 Gly Arg Glu Thr Val Ile Glu Tyr Leu Val Ser Phe Gly Val Trp Ile

353 130 135 140

356 Arg Thr Pro Pro Ala Tyr Arg Pro Pro Asn Ala Pro Ile Leu Ser Thr

357 145 150 155 160

360 Leu Pro Ala Trp Ala Arg Val Ile Asn

361 165

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367 <213> ORGANISM: Artificial Sequence

369 <220> FEATURE:

370 <223> OTHER INFORMATION: Synthetic

372 <400> SEQUENCE: 19

373 cacaaggacc tcttgagca gaagcgggcc gcggtggaca cctactgcgt agat

54

376 <210> SEQ ID NO: 20

377 <211> LENGTH: 54

378 <212> TYPE: DNA

379 <213> ORGANISM: Artificial Sequence

381 <220> FEATURE:

382 <223> OTHER INFORMATION: Synthetic

384 <400> SEQUENCE: 20

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54

388 <210> SEQ ID NO: 21

389 <211> LENGTH: 5

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/786,774

DATE: 07/21/2004
TIME: 09:27:26

Input Set : A:\08550~1.txt

Output Set: N:\CRF4\07212004\J786774.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 2,3,4 ✓
Seq#:4; Xaa Pos. 3,4 ✓
Seq#:21; Xaa Pos. 2,3,4
Seq#:25; Xaa Pos. 3,4
Seq#:26; Xaa Pos. 3,4
Seq#:27; Xaa Pos. 3,4
Seq#:30; Xaa Pos. 1,7
Seq#:31; Xaa Pos. 1,7
Seq#:32; Xaa Pos. 1,7
Seq#:33; Xaa Pos. 1,7
Seq#:34; Xaa Pos. 1,7
Seq#:35; Xaa Pos. 1,7
Seq#:36; Xaa Pos. 1,7

VERIFICATION SUMMARY

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Input Set : A:\08550~1.txt

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:410 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:478 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:504 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:530 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:684 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:706 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:750 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
L:772 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:794 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:816 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0